

- 1 -

piece 1, NC_000913, ynjH_gdhA-, config: linear, direction: -, begin: 1840424, end: 1840140

The figure shows a protein sequence alignment and a phylogenetic tree for the gene NC_000913.gdhA. The top part displays the amino acid sequence with various mutations highlighted by colored dots (red, green, blue) above the sequence. The bottom part shows a phylogenetic tree with three main clades, each containing multiple sequences. The tree is rooted at the bottom left and has a scale bar of 1.9 bits.

p35 1.9 bits
 p10 4.6 bits
 } p35-(23)-p10 1840357 Gap 1.4 bi
 p35-p10 1840357 total 5.0 bits

p35 5.2 bits

p10 2.8 bits

ir ynjH gdhA-

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{ } p35-(23)-p10 1840311  
----- p35-p10 1840311 total
```

ap 1.4 bits {-----} sd-(7)-ir 1840285 Gap 3.7 bits
6.5 bits -----| sd-ir 1840285 vnjH gdhA- total 6.8 bits

5' *1840180 *1840170 *1840160 *1840150 *1840140 3'

ir ynjH_gdhA-

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{-----} sd-(9)-ir 1840159 Gap 2.3 bits  
{-----} sd-ir 1840159 vniH gdhA- total 8.6 bits
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